

Figure 1. Amino Acid Sequences of Murine 1A6 and Human Consensus Sequences of Heavy Chain Subgroup III (HumIII) and Light Chain κ Subgroup I (Hum κ I).

V_H Domain

	1	11	21	31	41	
Mouse1A6	EVQLQ	QSGAE	LVKPG	ASLKL	SCTAS	GFNIK
					DTYIH	WMKQR
					PEQGLEW	I GR
		** **	* *	*		** *

HumIII	EVQLVESGGG	LVQPGGSLRL	SCAASGFNFS	-----WVRQA	PGKGLEWVA	---
--------	------------	------------	------------	------------	-----------	-----

	51 a	61	71	81 a bc	91	
Mouse1A6	IDP	PANDNTIYD	PKVQ	GKATMT	ADTSS	NTAYL
					QLNSLTSEDTAVY	YCTT
			** *	*		**

HumIII	-----A	DSVKGRF	T IS	RDDSKNTAYL	QMNSLRAEDTAVY	YCTT
--------	--------	----------------	------	-------------------	----------------------	-------------

		103	111	
Mouse1A6	SGYWFA	YWQG	TLVT	VSS

HumIII	-----	WGQG	TLVT	VSS
--------	-------	-------------	-------------	------------

V_L Domain

	1	11	21	31	41	51
Mouse1A6	DIVLTQSPAT	LSVTPGDSVS	LSCRASQ	SIS	NNLHWYQ	QKH
					SESPRLLIK	H ASQ
	**	**	***	* **		*

Hum κ I	DIQMTQSPSS	LSASVGDRVT	ITC-----	-----WYQQKP	GKAPKLLIY	-- ----
----------------	------------	------------	----------	-------------	-----------	---------

	61	71	81	91	101	
Mouse1A6	SISG	I PS	RFSGSGSGTD	FTLSINSVET	EDFGMFFCQQ	SNSWPYTFGG
						GTKLEIKR
	*		* **	***		*

Hum κ I	---GVPS	RFSGSGSGTD	FTLTSSLQP	EDFATYYC	----	-----FGQ
						GTKVEIKR

The CDR residues as defined by both Kabat and Chothia are shown in boldface.

Fig. 2A

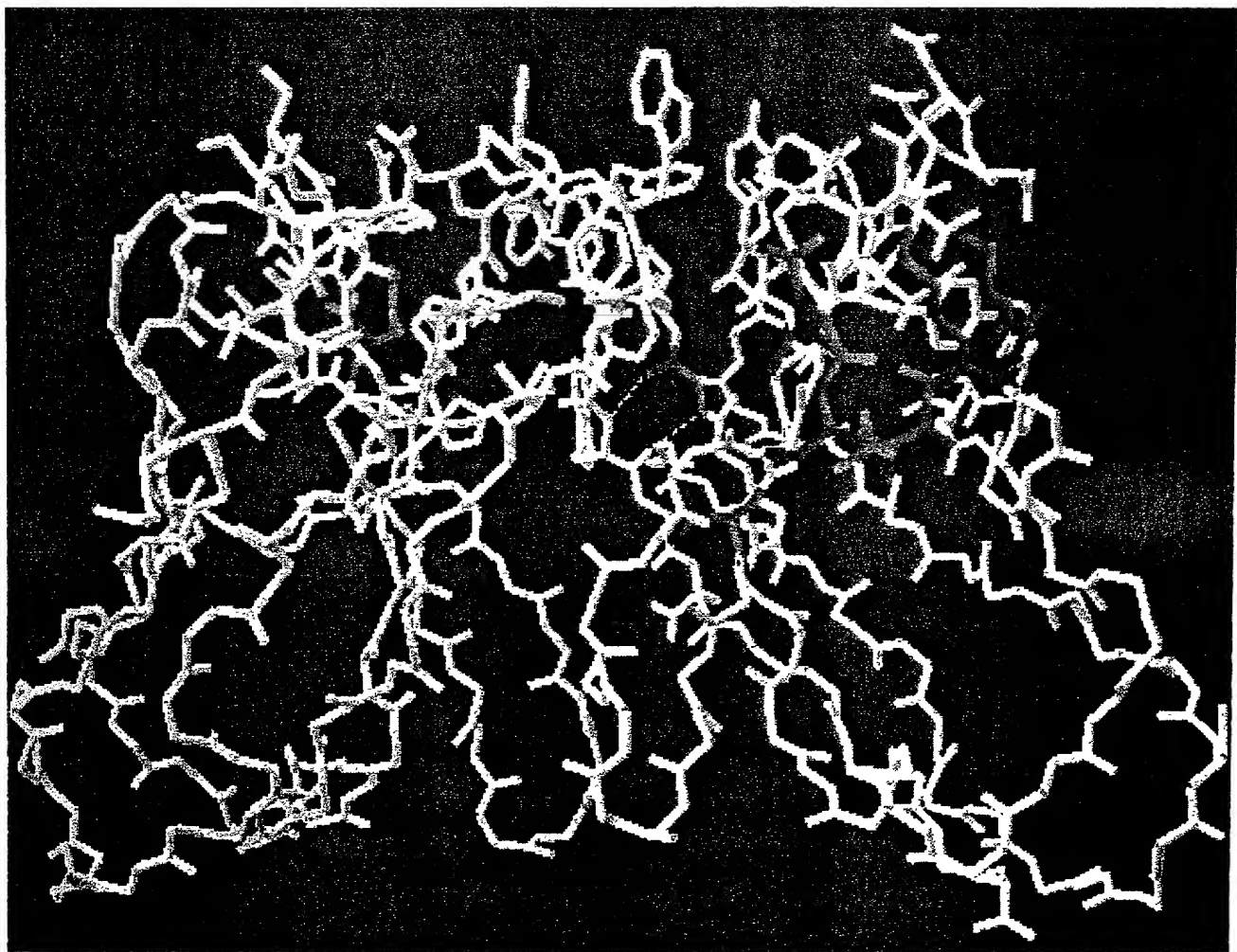


Fig. 2B

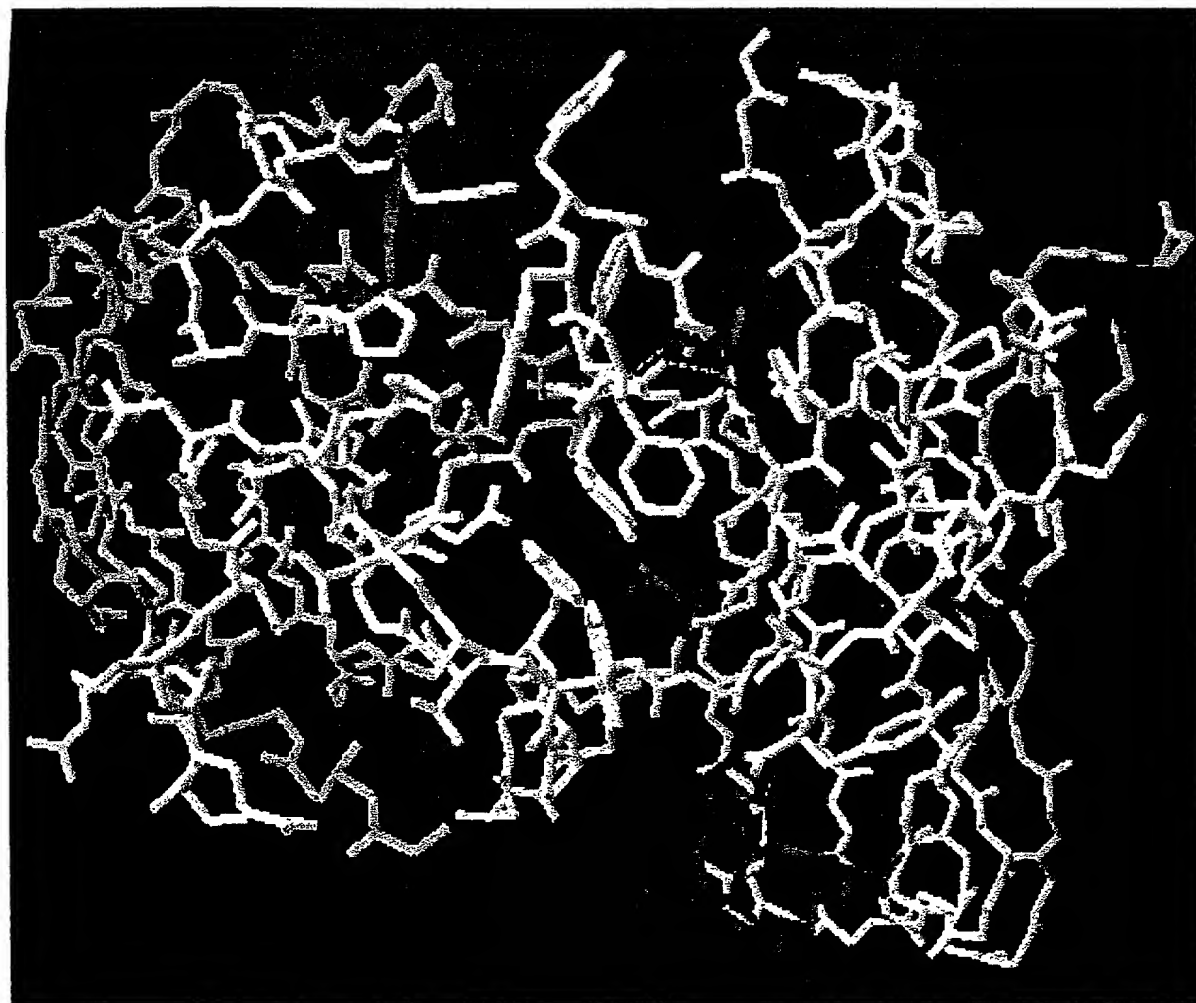


Fig. 2C

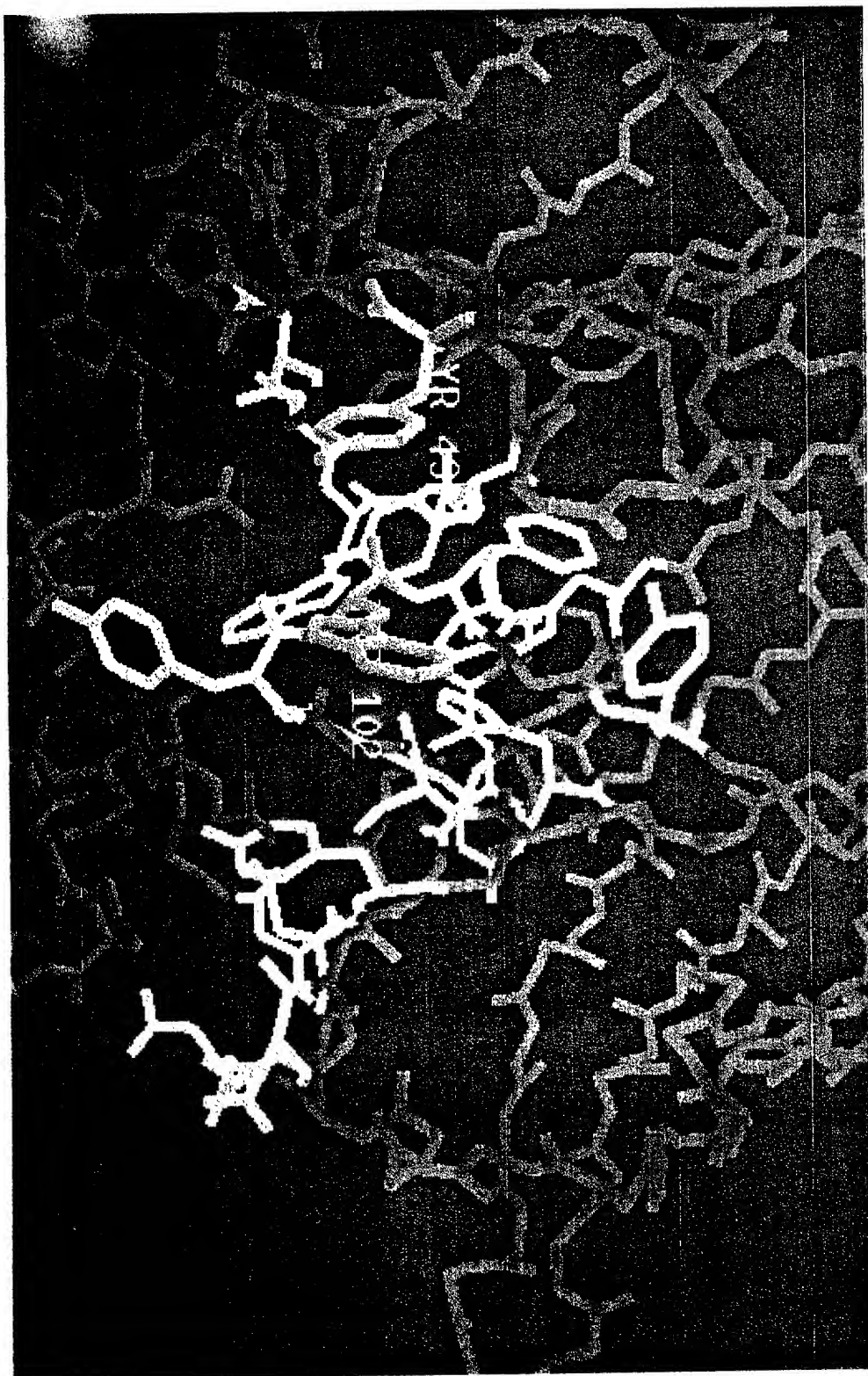


Figure 3. Amino Acid Sequences of Murine 1A6, Humanized 1A6 (Hum19), and Human Consensus Sequences of Heavy Chain Subgroup III (HumIII) and Light Chain κ Subgroup I (Hum κ I).

V_H Domain	
	1 11 21 31 41
Mouse	EVQLQQSGAE LVKPGASLKL SCTASGFNIK DTYIHW MKQR PEQGLEWI GR * * * * *
Hum19	EVQLVESGGG LVQPGGSLRL SCAASGFNIK DTYIHW VQRQA PGKGLEWVAR
HumIII	EVQLVESGGG LVQPGGSLRL SCAASGFNFS -----WVRQA PGKGLEWVA—
	51 a 61 71 81 abc 91
Mouse	IDPANDNTIYD PKVQGKATMT ADTSS NTAYL QL NSLTSEDTAVY YCT T * * * * *
Hum19	IDPANDNTIYA DSVKG RFT IS SDDSKNTAYL QMNSLRAEDTAVY YCTA * * *
HumIII	-----A DSVKG RFT IS RDDSKNTAYL QMNSLRAEDTAVY YCTR
	103 111
Mouse	SGYWFA YWGQGTLVT VSS
Hum19	SGYWFA YWGQGTLVT VSS
HumIII	-----WGQGTLVT VSS
V_L Domain	
	1 11 21 31 41 51
Mouse	DIVLTQSPAT LSVTPGDSVS LSCRASQ ^S IS NNLHWYQ ^Q KH SESPRLLIKH ASQ * * * * *
Hum19	DIQMTQSPSS LSASVGDRVT ITCRASQ ^S IS NNLHWYQ ^Q KP GKAPKLLIYH ASQ
Hum κ I	DIQMTQSPSS LSASVGDRVT ITC----- WYQ ^Q KP GKAPKLLIY -- ----
	61 71 81 91 101
Mouse	SISG I PS RFSGSGSGTD FTL ^S INSVET EDFGMFFC ^Q Q SNSWPYTFGG GTKLEIKR * * * * *
Hum19	SISG VPS RFSGSGSGTD FTLTISSLQ ^P EDFATYYC ^Q Q SNSWPYTFGQ GTKVEIKR
Hum κ I	---GVPS RFSGSGSGTD FTLTISSLQ ^P EDFATYYC --- ----- FGQ GTKVEIKR

The CDR residues as defined by both Kabat and Chothia are shown in boldface.

Figure 4. cDNA Sequences of Humanized scFv3 (Hum3) [SEQ ID. 2].

The restriction sites are underlined. CCATGG NCO I SITE; GGATCC BAMH I SITE; GTTAAC HPA I SITE

CGAACCATGGGCGATATCcgatgACCCAATCTCCGtctagcCTGAGCgccAG
TgttGGTgatCGAGTTaccattactTGCCGCGCCAGCCAATCTATCAGTAATAATCTTC
ACTGGTATCAACAAaaaccgggtaaagctCCGaaaCTTCTTATCAAACACGCCTCTCAG
AGCATTAGCGGCggtCCGAGCCGCTTCTCTGGCTCTGGCTCGGGCACGGACTTT
ACCCTTaccATCAGCTCTcttcagccgGAAGACtttGCCaccTATtatTGTCAGCAGTCTAA
TAGCTGGCCGTATACCTTCGGTcaaGGTACCAAGgtcGAGATTAAGCGCGGCGG
TGGCGGTTCTGGTGGCggtgtagcggtaggcGGTGGATCCGGTGGCGGTGGCAGCGA
AGTTCAACTTGTTGAGTCTGGTGGCGGTCTGGTTCAGCCGGGTGGCTCTCTGC
GCCTGTCTTGCGCAGCAAGCGGTTTCAACATTAAGGACACCTACATCCATTGG
atgAGGCAAGCTCCGGGTAAGGGTCTGGAGTGGGTGGCACGTATCGACCCGGC
AAACGACAACACCATTTACGATCCGAAGGTGCAGGGCCGTTTTACTatgTCTGC
GGACacCTCTAAGAACACCGCGTACCTTCAGATGAACTCTCTGCGTGCCGAGG
ACACCGCCGTCTACTACTGCACGACCTCTGGCTACTGGTTTGCCTACTGGGGC
CAGGGCACGCTTGTCACCGTCTCTTCTGGTTaACCC

Figure 5. Protection of HRV15 infection by mouse 1A6 scFv (Ms1) and humanized 1A6 scFv proteins (Hs3, 4, 7, 17, 18, 19 and 21).

